

AMENDMENTS TO THE CLAIMS:

Claim 16 is amended. The following is the status of the claims of the above-captioned application, as amended.

Claims 1-13 (Canceled).

Claim 14 (Previously presented). A composition comprising at least two thermostable enzymes selected from the group consisting of endoglucanase, xylanase, phytase, protease, galactanase, mannanase, dextranase, and alpha-galactosidase, wherein each of the thermostable enzymes has a melting temperature, T_m , of at least 70°C, as determined by Differential Scanning Calorimetry (DSC) at a pH in the interval of 5.0 to 7.0.

Claim 15 (Previously presented). The composition of claim 14, which comprises the following thermostable enzymes: (i) endoglucanase and xylanase; (ii) endoglucanase and protease; (iii) endoglucanase, xylanase and phytase; (iv) endoglucanase, xylanase and protease; (v) endoglucanase, xylanase, phytase and protease; (vi) xylanase and phytase; (vii) xylanase and protease; (viii) phytase and protease; (ix) phytase, protease and galactanase; (x) xylanase, phytase and protease; (xi) xylanase, protease and galactanase; (xii) phytase and galactanase; (xiii) galactanase and protease; (xiv) phytase, galactanase and alpha-galactosidase; (xv) phytase and alpha-galactosidase; (xvi) protease and alpha-galactosidase; (xvii) galactanase and alpha-galactosidase; (xviii) galactanase, protease and alpha-galactosidase; or (xix) at least two of endoglucanase, xylanase, phytase and galactanase.

Claim 16 (Currently amended). The composition of claim 14, which comprises

(A) at least one polypeptide having xylanase activity, which is a family 11 glycoside hydrolase; and

(B) at least one polypeptide having endoglucanase activity, which

(i) has an amino acid sequence of at least 75% identity to amino acids 4 -30 to 335 305 or 34 1 to 335 305 of SEQ ID NO:2 or 1 to 303 of SEQ ID NO: 18,

(ii) is encoded by a nucleic acid sequence which hybridizes under low stringency conditions with

(a) the mature endoglucanase encoding part of the plasmid contained in *Escherichia coli* DSM 14541,

(b) nucleotides 1 to 1008 or ~~90~~ 91 to 1008 of SEQ ID NO:1 or 97 to 1008 of SEQ ID NO: 17,

(c) a subsequence of (a) or (b) of at least 100 nucleotides, or

(d) a complementary strand of (a), (b) or (c);

(iii) is a fragment of (i) or (ii) that has endoglucanase activity.

Claim 17 (Previously presented). The composition of claim 16, wherein

(a) the endoglucanase and the xylanase are thermostable;

(b) the polypeptide having endoglucanase activity is a family 5 glycoside hydrolase;
and/or

(c) the polypeptide having xylanase activity is derived from a strain of *Aspergillus*, *Bacillus*, *Humicola*, *Thermomyces*, or *Trichoderma*.

Claim 18 (Previously presented). The composition of claim 14, further comprising

(a) at least one fat soluble vitamin, and/or

(b) at least one water soluble vitamin, and/or

(c) at least one trace mineral.

Claim 19 (Previously presented). The composition of claim 14 which is an animal feed additive.

Claim 20 (Previously presented). An animal feed composition having a crude protein content of 50 to 800 g/kg and comprising the composition of claim 14.

Claim 21 (Previously presented). The animal feed composition of claim 20, which comprises at least one of soy, wheat, barley, oats or rye.

Claim 22 (Previously presented). A method for the treatment of vegetable proteins, comprising the step of adding the composition of claim 14 to at least one vegetable protein or protein source.

Claim 23 (Previously presented). The method of claim 22, wherein the vegetable protein source comprises at least one of soy, wheat, barley, oats and rye.

Claim 24 (Previously presented). A method for improving the nutritional value of an animal feed, comprising adding the composition of claim 14 to the feed.